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	DB=PG	PB,USPT,EPAB,JPAB,DWPI; PLUR=YES; OP=OR	
Γ.	L56	L55 and @pd > 20060324	0
<u> </u>	L55	L50 and detect\$3	27
	L54	L50 and (biotin\$ or (biotin near2 "dATP"))	19
Γ	L53	L50 same (biotin\$ or (biotin near2 "dATP"))	0
Γ	L52	L50 same (biotin\$ or biotin near2 "dATP")	0
Γ.	L51	uracil same (endonuclease near2 "IV") same (nick\$2 or gap\$3 or abasic)	34
Γ.	L50	(displac\$4 near (nick\$2 or gap\$3 or abasic)) same label\$4	27
Г	L49	L48 and @pd > 20060412	0
	L48	phi29 near20 (reduce\$ or decrease\$ or mutat\$2) near5 (exonuclease)	. 6
Г	L47	phi29 near20 (exonuclease)	53
Г	L46	phi29 near (exonuclease)	1
Γ.	L45	phi29 near (reduce\$ or decrease\$)	. 0
<u>. </u>	L44	phi29	310
	DB = US	PT; PLUR=YES; OP=OR	
<u> </u>	L43	6762022.pn.	1
	DB=PG	PB,USPT,EPAB,JPAB,DWPI; PLUR=YES; OP=OR	
r.	L42	L40 same (label or biotin) and (array or microarray or chip or biochip)	58
L	L41	L40 same (label or biotin) same (array or microarray or chip or biochip)	2
Γ	L40	(cDNA) near ((oligo near2 dT) or (random) or (primer))	3855
Γ	L39	((cDNA) near ((oligo near2 dT) or (random) or (primer)) near50 biotin)	13
Γ	L38	((cDNA) near ((oligo near2 dT) or (random) or (primer)) near20 biotin)	10
Г	. L37	((cDNA) near ((oligo near2 dT) or (random) or (primer)) near15 biotin)	10
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Γ.	L36	•	1
	DB=PG	PB; PLUR = YES; OP = OR	
Γ.	L35	L34 and @pd > 20060412	13
	L34	McGall.in.	72
		PT; PLUR=YES; OP=OR	
Γ	L33	Mcgall.in.	72
	L32	L28 and nucleic	119
		PPB; PLUR=YES; OP=OR	0.5
Γ	L31	L28 and nucleic	86
	DB=US	PT; PLUR=YES; OP=OR	•

Γ	L30	L29	•	·	119	
	DB=PC	GPB, USPT; PLUR=YES; OP=OR				
Γ	L29	L28 and nucleic			205	
Γ	L28	Cole.in.			6222	
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Γ.	L27	20040166493			1	
Γ	L26	20040005614			1	
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	L25	5683896.pn.			1	
<u> </u>	L24	5536649.pn.			1	
	DB=PC	GPB; PLUR=YES; OP=OR		·		
Γ	L23	2005026147			0	
Γ	L22	Walker.in.			2853	
Γ	L21	Porat.in.			67	
Γ	L20	2004166493		•	0	
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	L19	6858413.pn.			1	
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Γ	L17	6197557.pn.			1	
Γ	L16	L15 and @pd > 20060317		•	0	
Γ	L15	5683896.pn.			1	
Γ	L14	6518026.pn.	·		1	
Γ.	L13	5536649.pn.			1	
Γ	L12.	5648211.pn.			1.	
Γ	L11	6117634.pn.			. 1	
Г	L10	6284460.pn.			1	
Γ.	L9	5858659.pn.			1	
Γ.	L8	6300063.pn.			1	
Γ	L7	63000063.pn.			0	
Γ	L6	5856092.pn.		•	1 '	
Γ	L5	6582938.pn.		·	. 1	
r.	L4.	6482804.pn.			1	
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Г	L3	20040067559			1	
Γ	Ľ2	20050136417			1	
Г	L1	20050026147			1	



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2.	Gene expression microarray analysis in cancer biology, pharmacology, and drug development: progress and potential • DISCUSSION Biochemical Pharmacology, Volume 62, Issue 10, 15 December 2001, Pages 1311-1336 Paul A. Clarke, Robert te Poele, Richard Wooster and Paul Workman SummaryPlus Full Text + Links PDF (459 K) Plant gene expression profiling with DNA microarrays • REVIEW ARTICLE					
	Plant Physiology and Biochemistry, Volume 39, Issue 11, November 2001, Pages 917-926 Shu-Hsing Wu, Katrina Ramonell, Jeremy Gollub and Shauna Somerville SummaryPlus Full Text + Links PDF (472 K)					
3. 「	[1] Preparation of cDNA from single cells and subcellular regions • ARTICLE Methods in Enzymology, Volume 303, 1999, Pages 3-18 Janet Estee Kacharmina, Peter B. Crino and James Eberwine Abstract Abstract + References PDF (3085 K)					
a 4 c	Identification of low-abundance differentially expressed transcripts using arrayed cDNA clones • ARTICLE Comparative Biochemistry and Physiology Part B: Biochemistry and Molecular Biology, Volume 133, Issue 4, December 2002, Pages 537-542					
***************************************	P. Golby, S. K. Stephens, J. P. Rast and J. F. Burke SummaryPlus Full Text + Links PDF (406 K)					
5 . 厂	Chronic neuropathic pain is accompanied by global changes in gene expression and shares pathobiology with neurodegenerative diseases • ARTICLE Neuroscience, Volume 114, Issue 3, 11 October 2002, Pages 529-546 H. Wang, H. Sun, K. Della Penna, R. J. Benz, J. Xu, D. L. Gerhold, D. J. Holder and K. S. Koblan SummaryPlus Full Text + Links PDF (937 K)					



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:15,418 total | 1,509 journal results | 13,392 preferred web results | 517 other web results

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1. Methods of amplifying sense strand RNA

Xu, Zhidong / Jablons, David / You, Liang / He, Biao, UNITED STATES PATENT AND TRADEMARK OFFICE PRE-GRANT PUBLICATION, Jun 2003

Email checked results

...sequence. Once cDNA synthesis is...to generate sense strand mRNA from the mRNA/cDNA heteroduplex...translation for second strand cDNA synthesis, and...the antisense cDNA strand, which...template" method, sense strand mRNA is generated...

Full text available at patent office. For more in-depth searching go to CexisNexis view all 13281 results from Patent Offices

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2. METHOD OF AMPLIFYING MRNA AND CDNA IN MICROQUANTITIES

TAKIGUCHI, Masaki, EUROPEAN PATENT APPLICATION, Nov 2003

...a double-stranded cDNA by using a sense strand cDNA in supernatant as a template, and...synthesizing an antisense strand cDNA and a sense strand cDNA on a carrier, (3) a process of adding...double-stranded cDNA by using said sense strand cDNA dissociated herein as a template and...

Full text available at patent office. For more in-depth searching go to LexisNexis

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3. IMMOBILIZED CDNA LIBRARIES

OTA, Toshio / MITSUHASHI, Masato / ISOGAI, Takao / WAKAMATSU, Ai,

EUROPEAN PATENT APPLICATION, Jul 2001

...terminal of a first strand cDNA can be any sequence...complementary sequence. A sense strand cDNA is synthesized by priming...At this time, the second strand cDNA synthesized is immobilized...used, a synthesized sense strand cDNA library finally...

Full text available at patent office. For more in-depth searching go to 🍎 LexisNexis view all 13281 results from Patent Offices

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4. METHOD OF AMPLIFYING MRNA AND CDNA IN MICROQUANTITIES

(TAKIGUCHI, Masaki) / Chiba, PATENT COOPERATION TREATY APPLICATION, Aug

...eliminated. By using the sense strand cDNA in the supernatant as a...to thereby amplify the cDNA mixture. By using the cDNA mixture, sense strand/antisense strand cRNA is...

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